SEQUENCE LISTING

(1) GENERAL INFORMATION:

- (i) APPLICANT: Sastry, Jagannadha K. Arlinghaus, Ralph B. Platsoucas, Chris D. Nehete, Pramod N.
- (ii) TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR ELICITING AN IMMUNE RESPONSE
- (iii) NUMBER OF SEQUENCES: 46
- (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: Arnold, White & Durkee
 - (B) STREET: P.O. Box 4433
 - (C) CITY: Houston
 - (D) STATE: Texas
 - (E) COUNTRY: USA
 - (F) ZIP: 77210
- (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: IBM PC compatible
 - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
- (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER: US 08/869,386
 - (B) FILING DATE: 05-JUN-1997
 - (C) CLASSIFICATION: Unknown
- (vii) PRIOR APPLICATION DATA:
 - -(A)=APPLICATION NUMBER: US 07/945,865
 - (B) FILING DATE: 16-SEP-1992
- (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Highlander, Steven L.
 - (B) REGISTRATION NUMBER: 37,642
 - (C) REFERENCE/DOCKET NUMBER: UTXC:538
 - (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: (512) 418-3000
 - (B) TELEFAX: (512) 474-7577
- (2) INFORMATION FOR SEQ ID NO:1:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:

- (XI) SEQUENCE DESCRIPTION: SEQ ID NO:1:

 Arg Ile Gln Arg Gly Pro Gly Arg Ala Phe Val Thr Ile Gly Lys
 1 5 10 15
- (2) INFORMATION FOR SEQ ID NO:2:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 19 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

Cys Arg Ile Lys Gln Ile Ile Asn Met Trp Gln Gly Val Gly Lys Ala 1 5 10 15

Met Tyr Ala

- (2) INFORMATION FOR SEQ ID NO:3:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 24 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

Asn Asn Thr Arg Lys Ser Glu Arg Ile Gln Arg Gly Pro Gly Arg Ala

1 10 15

Phe Val Thr Ile Gly Lys Ile Gly

- (2) INFORMATION FOR SEQ ID NO:4:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 13 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

The second secon

Glu Gln Leu Trp Val Thr Val Tyr Tyr Gly Val Pro Val

5 10

- (2) INFORMATION FOR SEQ ID NO:5:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 8 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

Arg Ala Phe Val Thr Ile Gly Lys
1 5

- (2) INFORMATION FOR SEQ ID NO:6:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 13 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

Thr Lys Gly Pro Gly Arg Val Ile Tyr Ala Thr Gly Gln
1 5 10

- (2) INFORMATION FOR SEQ ID NO:7:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 13 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:

His Ile Gly Pro Gly Arg Ala Phe Tyr Thr Thr Lys Asn 1 5 10

- (2) INFORMATION FOR SEQ ID NO:8:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 11 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:

Thr Tyr Gln Arg Thr Arg Ala Leu Val Thr Gly
1 5 10

- (2) INFORMATION FOR SEQ ID NO:9:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 16 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:

His Gly Glu Phe Ala Pro Gly Asn Tyr Phe Ala Leu Trp Ser Tyr Ala 1 5 10 15

- (2) INFORMATION FOR SEQ ID NO:10:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 25 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:

Cys Thr Arg Pro Asn Asn Asn Thr Arg Lys Ser Ile Arg Ile Gln Arg 1 5 10 15

Gly Pro Gly Arg Ala Phe Val Thr Ile 20 25

- (2) INFORMATION FOR SEQ ID NO:11:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 24 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:

Tyr Asn Lys Arg Lys Arg Ile His Ile Gly Pro Gly Arg Ala Phe Tyr

1 10 15

Thr Thr Lys Asn Asn Ile Gly Cys 20

(2) INFORMATION FOR SEQ ID NO:12:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:

Arg Ile His Ile Gly Pro Gly Arg Ala Phe Tyr Thr Thr Lys Asn
1 5 10 15

- (2) INFORMATION FOR SEQ ID NO:13:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:

Ser Leu Ser Ile Gly Pro Gly Arg Ala Pro Arg Thr Arg Glu Ile 1 5 10 15

- (2) INFORMATION FOR SEQ ID NO:14:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 24 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:14:

Asn Asn Thr Arg Lys Ser Ile Thr Lys Gly Pro Gly Arg Val Ile Tyr_
1 5 10 15

Ala Thr Gly Gln Ile Ile Gly Asp 20

- (2) INFORMATION FOR SEQ ID NO:15:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:15:
 - Gly Ile Ala Ile Gly Pro Gly Arg Thr Leu Tyr Ala Arg Glu Lys

(2)	INFORMATION	FOR	SEQ	ID	NO:16:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:16:

Ser Ile Thr Lys Gly Pro Gly Arg Val Ile Tyr Ala Thr Gly Gln
1 5 10 15

10

- (2) INFORMATION FOR SEQ ID NO:17:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:17:

Arg Val Thr Leu Gly Pro Gly Arg Val Trp Tyr Thr Thr Gly Glu

1 10 15

- (2) INFORMATION FOR SEQ ID NO:18:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:18:

Ser Ile His Ile Gly Pro Gly Arg Ala Phe Tyr Ala Thr Gly Asp 1 5 10 15

- (2) INFORMATION FOR SEQ ID NO:19:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:19:

INFORMATION FOR ORD IN 120 A													
INFORMATION FOR SEQ ID NO:20:													
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 15 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: (D) TOPOLOGY: linear 													
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:20:													
Ser Ile Tyr Ile Gly Pro Gly Arg Ala Phe His Thr Thr Gly Arg 1 5 10 15													
INFORMATION FOR SEQ ID NO:21:													
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 15 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: (D) TOPOLOGY: linear 													
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:21:													
Gly Ile His Phe Gly Pro Gly Gln Ala Leu Tyr Thr Thr Gly Ile 1 5 10 15													
INFORMATION FOR SEQ ID NO:22:													
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 15 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: (D) TOPOLOGY: linear													
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:22:													
Ser Ile Ser Ile Gly Pro Gly Arg Ala Phe Phe Ala Thr Thr Asp 1 5 10 15													
INFORMATION FOR SEQ ID NO:23:													
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 15 amino acids (B) TYPE: amino acid													

Ser Ile Arg Ile Gly Pro Gly Lys Val Phe Thr Ala Lys Gly Gly

(2)

(2)

(2)

(2)

(C) STRANDEDNESS:
(D) TOPOLOGY: linear

	Ser 1	Thr Pro Ile Gly Leu Gly Gln Ala	Leu 10	Tyr	Thr	Thr	Arg	Gly 15						
(2)	INFO	RMATION FOR SEQ ID NO:24:												
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 15 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: (D) TOPOLOGY: linear												
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:24:													
	Ser 1	Thr Pro Ile Gly Leu Gly Gln Ala	Leu 10	Tyr	Thr	Thr	Arg	Ile 15						
(2)	INFO	RMATION FOR SEQ ID NO:25:												
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 15 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: (D) TOPOLOGY: linear												
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO	:25:											
	Arg 1	Thr Pro Thr Gly Leu Gly Gln Ser	Leu 10	Tyr	Thr	Thr	Arg	Ser 15						
(2)	INFO	RMATION FOR SEQ ID NO:26:	-	· -		, -	-		-	. ·				
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 15 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: (D) TOPOLOGY: linear								5.				
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO	:26:		•									
	Arg 1	Ile His Ile Gly Pro Gly Ala Arg	Phe 10	Leu	Thr	Thr	Lys	Asn 15						

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:23:

(2) INFORMATION FOR SEQ ID NO:27:

(i) SEQUENCE CHARACTERISTICS:

(B) TYPE: amino acid

(A) LENGTH: 15 amino acids

		(xi)	SEQU	ENCE	DESC	RIPTIC	N: SI	EQ II	ои с	:27:								
		Arg	Ile	His I	le Gl	ly Pro	Gly	Arg	Ala	Phe 10	Phe	Thr	Thr	Lys	Asn 15			
	(2)	INFOR	MATI	ON FO	R SE	O ID N	iO:28	:										
		(i) :	(A) (B) (C)	LENG TYPE STRA	TH: 1 : ami	ACTERI 11 ami ino ac NESS: : line	no ao											
		(xi)	SEQU	ENCE	DESCI	RIPTIC	N: SI	EQ II	ои с	:28:								
		Val' 1	Tyr	Tyr G	ly Va 5	al Pro	Val	Trp	Lys	Glu 10	Ala					•		
	(2)	INFOR	MATI	ON FO	R SEG	Z ID N	10:29	:					-					
		(i)	(A) (B) (C)	LENG TYPE STRA	TH: : : am: NDEDI	ACTERI 13 ami ino ac NESS: : line	no ao											
		(xi)	SEQU	ENCE	DESCI	RIPTIC	N: S	EQ II	ои о	:29:								
-	·	Gly 1	Val		al Ti	rp Lys	Glu	Ala	Thr	10	Leu	Phe	_	-				-
	(2)	INFOR	MATI	ON FO	R SE	ID I	 IO:30	:									z. .	
		(i)	(A) (B)	LENG	TH: :	ACTERI 11 ami ino ac NESS:	no a											

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(2) INFORMATION FOR SEQ ID NO:31:

(i) SEQUENCE CHARACTERISTICS:

(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:30:

Ala His Lys Val Trp Ala Thr His Ala Cys Val

(C) STRANDEDNESS:
(D) TOPOLOGY: linear

- (A) LENGTH: 12 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS:
- (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:31:

Cys Val Pro Thr Asn Pro Val Pro Gln Glu Val Val 1 5 10

- (2) INFORMATION FOR SEQ ID NO:32:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 11 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:32:

Val Leu Glu Asn Val Thr Glu Asn Phe Asn Met 1 5 10

- (2) INFORMATION FOR SEQ ID NO:33:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 11 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:33:

Asn Asn Met Val Glu Gln Met His Glu Asp Ile

1 5 10

- (2) INFORMATION FOR SEQ ID NO:34:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 13 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:34:

Glu Gln Met His Glu Asp Ile Ile Ser Leu Trp Asp Gln
1 10

(2) INFORMATION FOR SEQ ID NO:35:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 13 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:35:

Leu Trp Asp Gln Ser Leu Lys Pro Cys Val Lys Leu Thr 1 5 10

- (2) INFORMATION FOR SEQ ID NO:36:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:36:

Ser Leu Lys Pro Cys Val Lys Leu Thr Pro Leu Cys
1 5 10

- (2) INFORMATION FOR SEQ ID NO:37:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 13 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:37:

Ser Val Ile Thr Gln Ala Cys Ser Lys Val Ser Phe Glu
1 5 10

- (2) INFORMATION FOR SEQ ID NO:38:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 14 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:38:

Phe Glu Pro Ile Pro Ile His Tyr Cys Ala Phe Pro Gly Phe 1 5 10

- (2) INFORMATION FOR SEQ ID NO:39:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 11 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:39:

Lys Lys Phe Asn Gly Thr Gly Pro Cys Thr Asn 1 5 10

- (2) INFORMATION FOR SEQ ID NO:40:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 13 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:40:

Gly Thr Gly Pro Cys Thr Asn Val Ser Thr Val Gln Cys
1 5 10

- (2) INFORMATION FOR SEQ ID NO:41:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 14 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:41:

Val Gln Cys Thr His Gly Ile Arg Pro Val Val Ser Thr Gln
1 10

- (2) INFORMATION FOR SEQ ID NO:42:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 13 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:42:

Tyr Leu Arg Asp Gln Gln Leu Leu Gly Ile Trp Gly Cys

10

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 25 amino acids

5

- (B) TYPE: amino acid
- (C) STRANDEDNESS:
- (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:43:

Phe Leu Gly Phe Leu Gly Ala Ala Gly Ser Thr Met Gly Ala Ala Ser 1 10 15

Leu Thr Leu Thr Val Gln Ala Asn Gln 20 25

- (2) INFORMATION FOR SEQ ID NO:44:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 28 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:44:

Cys Arg Ile Lys Gln Ile Ile Asn Met Trp Gln Gly Val Gly Lys Ala 1 5 10 15

Met Tyr Ala Pro Pro Ile Gly Gly Gln Ile Arg Cys 20 25

- (2) INFORMATION FOR SEQ ID NO:45:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 36 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:45:

Cys Thr Arg Pro Asn Asn Asn Thr Arg Lys Ser Ile Arg Ile Gln Arg

1 10 15

Gly Pro Gly Arg Ala Phe Val Thr Ile Gly Lys Ile Gly Asn Met Arg 20 25 30

Gln Ala His Cys

- (2) INFORMATION FOR SEQ ID NO:46:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 16 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:46:

Lys Gln Ile Ile Asn Met Trp Gln Glu Val Gly Lys Ala Met Tyr Ala 1 5 10 15